## Steps for hypothesis testing

Step 0: Define problem Objective

Step 1: Specify the Null Hypothesis.

Step 2: Specify the Alternative Hypothesis.

Step 3: Set the Significance Level

Step 4: Calculate the Test Statistic and Corresponding P-Value.

Step 5: Drawing a Conclusion.

In [ ]:

## Step 0: Define problem Objective:

Exercising does not affect weight`

This statement is my Hypothesis. Let's call it Null hypothesisfor now. For now, it is the status quo as in we consider it to be true

## Step 1 and 2: Specify Null and Alternate Hypothesis:

H°: Exercising does not affect weight. Or equivalently  $\mu$  = 0

 $H^A$ : Exercise does reduce weight. Or equivalently  $\mu$ >0

We collect weight loss data for a sample of 30 people who regularly exercise for over 3 months.

- WeightLoss Sample Mean = 2 kg
- weightLoss Sample Mean = 2 kg
   Sample Standard Deviation = 1 kg

Does this prove that exercise does reduce weight? Infact, it sort of looks like that exercising does have its benefits as people who exercise have lost on an average 2 kgs.

#### Step 3: Set the Significance Level

Assuming that the null hypothesis is true, what is the probability of observing a sample mean of 2 kg or more extreme than 2 kg?

- Assuming we can calculate this If this probability value is meagre (lesser than a threshold value), we reject our null hypothesis. And otherwise, we fail to reject our null hypothesis. Why fail to reject and not accept?
- This probability value is actually the p-value. Simply, it is just the probability of observing what

In [ ]:

Out[2]: 0.0

## Step 4 Calculate the test statastics and corresponding p-value

In [1]: from scipy.stats import norm import numpy as np

In [2]: ### loc is the location parameter and mean is the normal distribution p=1-norm.cdf(x=2,loc=0,scale=1/np.sqrt(30)) p

As the p value(0.0)is less than significant level(0.05) therefore we have to reject the null hypothesis(Exercising does not affect weight.)

### Step 5: Drawing the Conclusion

As such, this is a very small probability p-value (less than the significance level of 0.05) for the mean of a sample to take a value of 2 or more.

And so we can reject our Null hypothesis. And we can call our results statistically significant as in they don't just occur due to mere chance

import numpy as np
import pandas as pd

In [4]: cars\_df=pd.read\_csv("cars93.csv")

In [5]: cars\_df.head()

Out[5]:

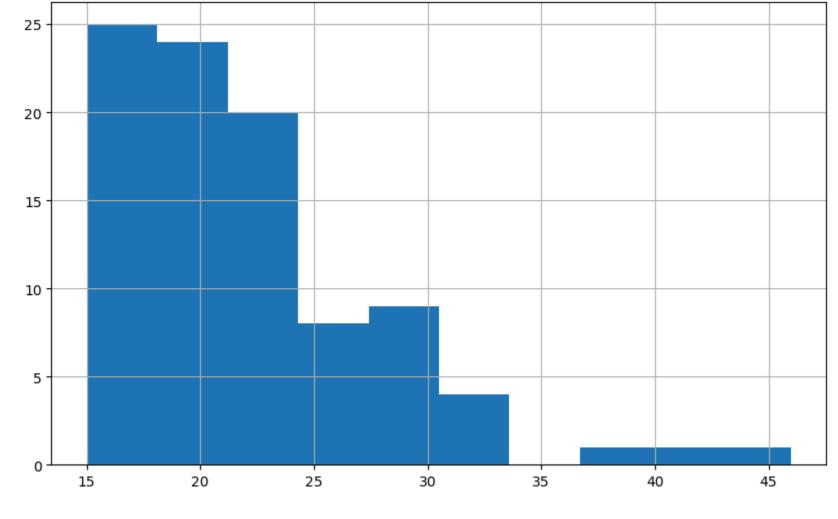
	Manufacturer	Model	Туре	Price	MPG.city	AirBags	Horsepower	Passengers	Rear.seat.room	Luggage.room
0	Acura	Integra	Small	15.9	25	Driver only	140	5	26.5	11.0
1	Acura	Legend	Midsize	33.9	18	Driver & Passenger	200	5	30.0	15.0
2	Audi	90	Compact	29.1	20	Driver only	172	5	28.0	14.0
3	Audi	100	Midsize	37.7	19	Driver & Passenger	172	6	31.0	17.0
4	BMW	535i	Midsize	30.0	22	Driver only	208	4	27.0	13.0

In [9]: cars\_df.shape

Out[9]: (93, 10)

In [6]: cars\_df["MPG.city"].hist(figsize=(10,6),bins=10)

Out[6]: <Axes: >



In above graph we can see that the data is right skweed

So for proving central limit theorem we will collect random sampels from the data and plot their mean

# In [7]: sampling\_distribution=[]

print(sampling\_distribution)

for i in range(500):
 #### generates random 50 integer index between 0 to len ("MPG.city")
 sample\_index=np.random.randint(0,len(cars\_df["MPG.city"]),50)
 ### Extract
 sample=cars\_df["MPG.city"][sample\_index]
 sample\_mean=sample.mean()
 sampling\_distribution.append(sample\_mean)

[21.44, 21.94, 23.58, 21.9, 22.38, 21.88, 22.7, 22.4, 21.96, 22.14, 22.14, 23.6, 23.68, 22.06, 21.46, 22.0, 21.6, 23.78, 21.02, 22.64, 22.94, 22.76, 23.16, 21.6, 22.02, 22.78, 23.08, 22.48, 23.0, 21.64, 21.18, 24.06, 21.58, 21.7, 22.4, 22.52, 21.96, 22.48, 21.36, 21.48, 22.92, 22.02, 21.96, 21.98, 22.66, 21.19, 22.94, 22.6, 21.6, 22.18, 23.8, 21.86, 22.22, 20.98, 21.86, 22.12, 21.90, 22.16, 21.90, 22.14, 21.46, 23.16, 22.28, 21.90, 22.14, 21.46, 23.16, 22.28, 21.90, 22.14, 21.46, 23.16, 22.28, 21.90, 22.18, 22.18, 21.68, 22.29, 21.90, 21.14, 21.56, 21.14, 21.56, 21.14, 21.56, 21.08, 23.4, 22.66, 21.44, 21.74, 22.8, 21.77, 21.66, 22.8, 21.90, 22.15, 23.94, 22.14, 21.90, 22.14, 23.02, 22.18, 21.90, 22.16, 22.18, 23.44, 22.18, 21.68, 22.14, 23.02, 22.18, 21.90, 22.14, 23.02, 22.08, 21.12, 21.90, 22.14, 23.02, 22.08, 21.12, 21.92, 22.7, 21.26, 23.5, 21.76, 22.06, 23.14, 22.18, 21.66, 21.82, 21.92, 22.74, 22.54, 22.22, 22.12, 21.92, 22.14, 23.02, 22.08, 21.12, 21.92, 22.74, 22.8, 22.8, 22.84, 21.84, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.18, 21.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 21.04, 22.14, 22.18, 21.14, 21.04, 22.18, 21.14, 21.04, 22.14, 21.04,

In this above cell we declared one empty dictionary ie sampling\_distribution which holds random samples, in for loop we are generating random numbers within range of 0 to length of our data and after we are taking the mean of those sample. After taking mean we are appending that value in our empty list

# In [8]: pd.Series(sampling\_distribution).hist(figsize=(10,6),bins=10)

